Phylogenetic position of the bivalve family Cyrenoididae removal from (and further dismantling of) the superfamily Lucinoidea

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ABSTRACT

A molecular analysis using sequences from 18S and 28S rRNA genes of the brackish and freshwater bivalve *Cyrenoida floridana*, in conjunction with a wide range of other heterodont bivalves, demonstrated a close relationship with the families Corbiculidae and Glauconomidae and distant from the Lucinoidea, where the Cyrenoididae had been usually classified. Based on this result it is proposed that the Cyrenoididae be removed from the Lucinoidea, which, for living taxa, now includes only the family Lucinidae.

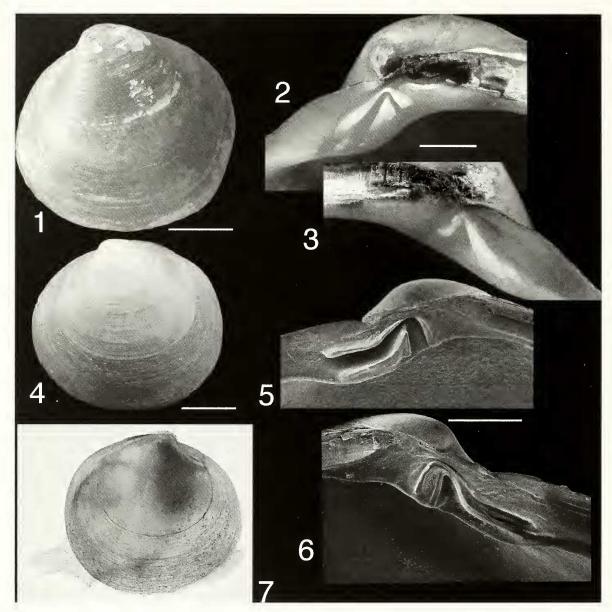
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INTRODUCTION

Until recently, most classifications of bivalve mollusks ineluded within the superfamily Lueinoidea several component families (Lueinidae, Fimbriidae, Thyasiridae, Ungulinidae Cyrenoididae, and fossil Mactromyidae) (e.g. Dall, 1901; Chavan, 1969; Boss, 1982, Vaught, 1989; Amler, 1999). The Lucinidae and some Thyasiridae are notable for the ehemosymbiosis with sulphideoxidizing baeteria housed in the ctenidia (Southward, 1986; Taylor and Glover, 2006). Molecular analyses of the Lueinoidea, eompared with a wide range of other heterodont bivalves, demonstrated that superfamily was not monophyletie, with the Thyasiridae and Ungulinidae not elosely related to the Lucinidae (Williams, Taylor, and Glover, 2004; Taylor, Williams, and Glover, 2007; Taylor et al., 2007). The Ungulinidae group near families such as the Veneridae, Corbiculidae, and Mactridae, while the Thyasiridae form a basal clade within the Euheterodonta and are eonsidered as a distinct superfamily—Thyasiroidea. Fimbria fimbriata, one of the two living species of Fimbriidae, nested together with Lucinidae species, with no support for separate familial status, and the nominal family was synonymized accordingly. Apart from the Lucinidae, the only other family with living species still elassified within Lucinoidea is the Cyrenoididae, but lack of suitably preserved material has precluded inelusion in molecular analyses. From morphological evidence, Williams et al. (2004) and Taylor and Glover (2006) thought a relationship to the Lucinidae unlikely, *Cyrenoida* having medium to long fused siphons, well developed labial palps and ctenidia with two demibranehs. These statements led Bieler and Mikkelsen (2006) to place Cyrenoididae as *incertae sedis*.

The Cyrenoididae Adams and Adams, 1857 (= Cyrenellidae Gray, 1853) eomprise a small group of around ten nominal speeies of little-studicd bivalves inhabiting brackish to freshwaters, classified into a single genus, *Cyrenoida* (type speeies *C. dupontia* Joannis, 1835) (Figures 1–3), distributed in western Africa, eastern and western Americas and some islands of the Caribbean. The West African speeies inhabit braekish mangrove habitats (Pilsbry and Bequaert, 1927). In the eastern USA, *Cyrenoida floridana* Dall, 1901 (Figures 4–7) ranges from Delaware to the eoast of the Gulf of Mexico, maybe as far west as Yueatan (Vokes and Vokes, 1983), where it inhabits fresh and brackish water habitats (Leathem, Kinner, and Maurer, 1976; Kat, 1982; Bishop and Haekney, 1987).

Dall (1895) was the first to place the Cyrenoididae within the Lueinoidea, stating (p. 545) "These are estuarine Lueinacea." Later (Dall, 1901: 817) stated "...shells of this group with a Lucinoid animal and *Diplodonta*like shell, exhibit a hinge structure which is wholly distinct from any other of the Lucinacea." Many later classifications, including the influential Treatise of Invertebrate Paleontology (Chavan, 1969), followed Dall in placing the Cyrenoididae within the Lucinoidea (e.g. Vokes, 1980; Boss, 1982, Vaught, 1989, Skelton and Benton, 1993; Amler, 1999).



Figures 1–7. Cyrenoida dupontia (1–3) and C. floridana (4–7). 1. Cyrenoida dupontia Joannis, left valve, Senegal (BMNH 20081055). Seale bar = 10 mm. 2–3. Hinge teeth of Cyrenoida dupontia, right valve (2) and left valve (3). Seale bar = 2 mm. 4. Cyrenoida floridana Dall, left valve, Blue Hole, Big Pine Key, Florida (BMNH 20081054). Seale bar = 2 mm. 5–6. Hinge teeth (SEM images) of Cyrenoida floridana, right valve (5) and left valve (6) (BMNH 20081054). Seale bar = 500 µm. 7. Cyrenoida floridana, living speeimen with short, fused siphons, Blue Hole, Big Pine Key, Florida. (Photo R. Bieler, September 2007).

Nevertheless, different opinions were expressed by other authors, Fischer (1887: 1096), for example, placed Cyrenoididae (as Cyrenellidae) into a suborder Conehacea, near to *Corbicula* and Ungulinidac but apart from the Lucinoidea. While Thiele (1934) included Cyrenoididae with other fresh and brackish water bivalves in the *stirps* Sphaeriacca but not positioned closely to Lucinoidea. The family was elevated to superfamily status by Olsson (1961: 227) but placed near to Lucinoidea, a deeision also followed by Keen (1971). The superfamily Cyrenoidoidea was also recognized by Nevesskaya et al. (1971) and placed along with Lueinoidea in the order Astartida. Alternatively, and rather bizarrely, Starobogatov (1992) placed Cyrenoidoidea within the infraorder Eryeinoinei along with Cyamioidea, Galeonmatoidea and Leptonoidea, all eontained within the order Lueiniformes.

Clearly, there exists much uncertainty concerning the phylogenetic position of Cyrenoididae amongst the heterodont bivalves but this has never been tested by either morphological or molecular analyses. In 2007, we obtained samples of *Cyrenoida floridana* suitable for molecular analysis and in this paper we present 18S and 28S rRNA sequences for the species that enable us to establish the phylogenetic position of the family in relation to a wide range of heterodont bivalve taxa previously analysed (Taylor et al., 2007) and specifically address the question of whether the Lucinidae and Cyrcnoididae form a monophyletic group.

MATERIALS AND METHODS

The sample of *Cyrenoida floridana*, preserved in 100% ethanol (BMNH 20081053), was collected (18 September 2007) from Blue Hole (24°42.4′ N, 81°22.8′ W) a freshwater pond on Big Pine Key, Monroe County, Florida Keys, Florida, USA, from shoreline mud up to 0.5 m depth among roots of marginal reeds. Other material from the same site is lodged at the Field Museum of Natural History (FMNH 314434; 317667).

For the molecular analysis, methods of DNA extraction, amplification and sequencing followed by sequence analysis and phylogenetic reconstruction are as described in Taylor et al. (2007). Sequences for *Cyrenoida floridana* were analysed together with the data set of heterodonts listed in Taylor et al. (2007, Table 1), with the addition of new 18S and 28S sequences for *Mya arenaria* Linnaeus, 1758 (family Myidae) from Gydnia, Poland. The new sequences for *Cyrenoida floridana* and *Mya arenaria* are lodged in GenBank (Accession numbers: *C. floridana* FM999789, FM999790; *M. arenaria* FM999791, 779792). Voucher specimens of both species are housed in the Department of Zoology, The Natural History Museum, London.

Phylogenies were constructed using Bayesian methods (MrBayes v3.1.2, Huelsenbeck and Ronquist, 2001) using a GTR+G+I model. The analysis for each data set was run for 3,500,000 generations, with a sample frequency of 100. Each analysis was run twice. The first 15,000 trees from each run were discarded so that the final consensus tree was based on the combination of accepted trees from each run (a total of 40,000 trees). Support for nodes was determined using posterior probabilities (PP, calculated by MrBayes).

RESULTS

The combined tree based on concatenated sequences from 18S rRNA and 28S rRNA genes is shown in Figure 8. The individual trees based on single genes are very similar in topology to those published previously (Taylor et al., 2007). In all analyses *Cyrenoida floridana* nests in a highly supported clade with *Corbicula fluminea* (Corbiculidae) and *Glaneonome virens* (Glauconomidae). This clade forms part of a major group of heterodonts named Neoheterodontei by Taylor et al. (2007). *Cyrenoida* is widely separated from both Thyasiridae and Lucinidae that appear in the more basal parts of the tree. The Ungulinidae, although also a member of the Neoheterodontei, form a separate clade distinct from *Cyrenoida*.

DISCUSSION

It would have been desirable to have included the type species of Cyrenoida, namely, Cyrenoida dupontia Joannis, 1835, from West Africa, in the molecular analysis but no suitably preserved material was available. Although a much smaller species, C. floridana is similar to C. dupontia in shell characters, notably the unusual hinge dentition, and we feel confident that they are members of the same group. Cyrenoida dupontia has three cardinal teeth in the right valve, the anterior of these is thin and elongate and the central tooth larger and slightly bifid (Figures 2–3) while the left valve has two cardinals, the posterior tooth smaller and bifid and the anterior tooth elongate. Lateral teeth are absent. The dentition of C. floridana is very similar (Figures 4-6) with three cardinal teeth in the right valve, the central being larger and two cardinal teeth in the left valve with the posterior tooth bifid and the anterior tooth elongate. We have also examined the gross anatomy of *Cyrenoida* rosea (d'Ailly, 1896) from Nigeria (National Museum of Wales specimen NMW.Z.2003.029.02041) and this has ctenidia with two demibranchs, with the inner demibranch larger, paired triangular labial palps, and fused medium-length posterior siphons. Cyrenoida floridana is similar, with small outer demibranchs, triangular labial palps and short fused posterior siphons, the inhalant with papillae (Figure 7). Despite the presence of siphons, there is no pallial sinus in any *Cyrenoida* species.

The main conclusion of this study, based on our results for Cyrenoida floridana, is that the Cyrenoididae should be removed from the Lucinoidea and classified close to, or possibly within, the Corbiculoidea. The status of Cyre*noida* in relation to Corbiculidae and Glauconomidae needs further analysis with a larger dataset of corbiculid species. For the present the family can be classified within a separate superfamily Cyrenoidoidca as proposed by Olsson (1961). Molecular evidence for a highly supported relationship between Corbiculidae and Glauconomidac was reported by Taylor et al. (2007) although the elongate shells with deep pallial sinus and long siphons of *Glaueonome* are less similar morphologically to Cyrenoididae and Corbiculidae. Species of Cyrenoididae and Corbiculidae occur in both brackish and freshwater habitats while Glauconomidac live intertidally among mangroves in environments of fluctuating salinity.

For living taxa, we consider that the superfamily Lucinoidea should now include only the family Lucinidae, with the families Thyasiridac, Ungulinidae and Cyrcnoididae excluded. The position of the entirely fossil families Mactromyidae, Ilionidae, and Paracyclidae is unresolved although the latter two embrace species with lucinid characters.

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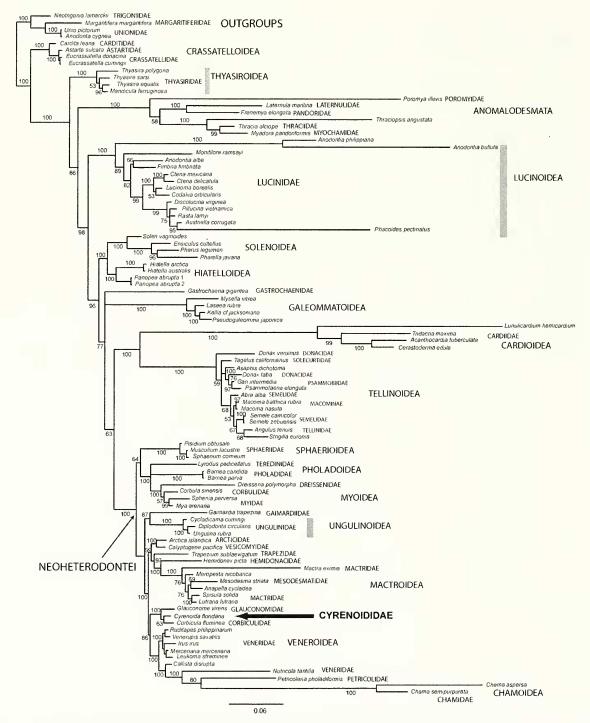


Figure 8. Molecular phylogeny of heterodont bivalves including *Cyrenoida* produced by Bayesian analysis for concatenated sequences from 18S and 28S rRNA genes. The tree was drawn using members of the palaeoheterodonts Trigoniidae, Unionidae, and Margaritiferidae as outgroups. Support values are posterior probabilities. Nodes with <50% support have been collapsed. Positions of Lucinoidea, Thyasiroidea, and Ungulinoidea marked by grey bars. Details of taxa in Taylor et al. (2007).

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